WHAT IS CLAIMED IS:

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1. An isolated and purified ATP diphosphohydrolase obtainable from bovine aorta characterized by the following physico-chemical properties:

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-a catalytic unit of a molecular weight on denaturing polyacrylamide gel electrophoresis of about 78 KDa in its native form;

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-a deglycosylated form of said catalytic unit of a molecular weight on SDS-PAGE of about 56 KDa; and characterized in that it comprises the amino acid sequences defined in SEQ. ID. NOs. 3 to 6.

2. An ATP diphosphohydrolase as defined in claim 1 further comprising the amino acid sequence defined in SEQ. ID. No.: 8 or a variant thereof.

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3. An isolated and purified ATP diphosphohydrolase obtainable from pig pancreatic zymogen granules

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characterized by the following physico-chemical properties:

-a catalytic unit of a molecular weight on denaturing polyacrylamide gel electrophoresis of about 54 KDa in its native form;

-a deglycosylated form of said catalytic unit of a molecular weight on SDS-PAGE of about 35 KDa; and characterized in that it comprises the amino acid sequence defined in SEQ. ID. NO.: 7.

- 4. A process for purifying an ATP-diphosphohydrolase enzyme from a tissue capable to convert ATP to ADP and ADP to AMP which comprises:
 - a) obtaining a sub-cellular microsomal fraction from an homogenate of said tissue;
- b) solubilizing said microsomal fraction in the presence of a non-ionic detergent;
 - c) centrifuging said solubilized microsomal fraction to obtain a supernatant containing said enzyme;

- d) submitting said supernatant to an ion-exchange chromatography to obtain a first enzyme eluate;
- e) submitting said first eluate to an affinity column chromatography to obtain a second enzyme eluate; and
- f) submitting said second eluate to a separation step on a non-denaturing gel electrophoresis to recover said enzyme free of any contaminant, the presence of said contaminant being monitored by overstaining said gel in a silver nitrate dye or Coomassie Blue dye.
- 5. A process according to claim 4 wherein said ion exchange chromatography is achieved on a column containing Diethylaminoethyl (DEAE).
- 6. A process according to claim 5 wherein said column is
 a DEAE agarose column.

- 7. A process according to claim 4 or 5 wherein said affinity column chromatography is achieved on an Affigel™ Blue column.
- 8. A process according to claim 4, 5, 6 or 7 wherein said non-ionic detergent is Triton $X-100^{\text{TM}}$.
- 9. A process according to claim 4, 5, 6, 7 or 8 wherein an aliquot of said enzyme is further submitted after step f) to a polyacrylamide gel electrophoresis under denaturing conditions to verify its homogeneity and to obtain its apparent molecular weight.
- 10. A process according to claim 9 wherein said enzyme is obtained from pig pancreatic zymogen granules and has an apparent molecular weight of 54 Kilodaltons.
- 11. A process according to claim 9 wherein said enzyme
 15 is obtained from bovine aortic intima layer and has an
 apparent molecular weight of about 78 Kilodaltons.

- 12. A process according to claim 10 wherein, between steps e) and f), a step of deglycosylation is included, and whereby the apparent molecular weight is shifted from 54 to 35 KDa.
- 13. A process according to claim 11 herein, between steps e) and f), a step of deglycosylation is included, and whereby the apparent molecular weight is shifted from 78 to 56 KDa.
- 14. The use of the ATP diphosphohydrolase of claim 1 or
 2, for reducing platelet aggregation and thrombogenicity.
 - 15. The use of an ATP diphosphohydrolase for reducing platelet aggregation and thrombogenicity, said ATP diphosphohydrolase having the amino acid sequence defined in SEQ. ID. NO.: 1, or a variant thereof, or a part thereof, said variant or part being capable of converting ATP to ADP and ADP to AMP.

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- 16. A composition for use in the reduction of platelet aggregation and thrombogenicity which comprises as an active ingredient the ATP diphosphohydrolase of claim 1 or 2 or an ATP diphosphohydrolase which sequence is defined in SEQ. ID. NO.: 1, or a variant or a part thereof, which variant orpart has an ATP diphosphohydrolase activity, in an acceptable pharmaceutical carrier.
- 17. A process for producing an ATP diphosphohydrolase which comprises the steps of:
- obtaining a host which comprises a nucleic acid encoding a protein having the amino acid sequence defined in SEQ. ID. NO.: 1, or a variant thereof, or a part thereof, said variant or part being capable of converting ATP to ADP and ADP to AMP;
- culturing said host in a culture medium supporting the growth of said host and the expression of said nucleic acid;

- recovering the ATP diphosphohydrolase from the culture medium or from said host; and
 - purifying the ATP diphosphohydrolase.
- 18. A process as defined in claim 17, wherein said nucleic acid has a sequence defined in SEQ. ID. NO.: 2, a variant thereof or a part thereof, said variant or part being capable of producing an ATP diphosphohydrolase which converts ATP to ADP and ADP to AMD.